

FIG. 1

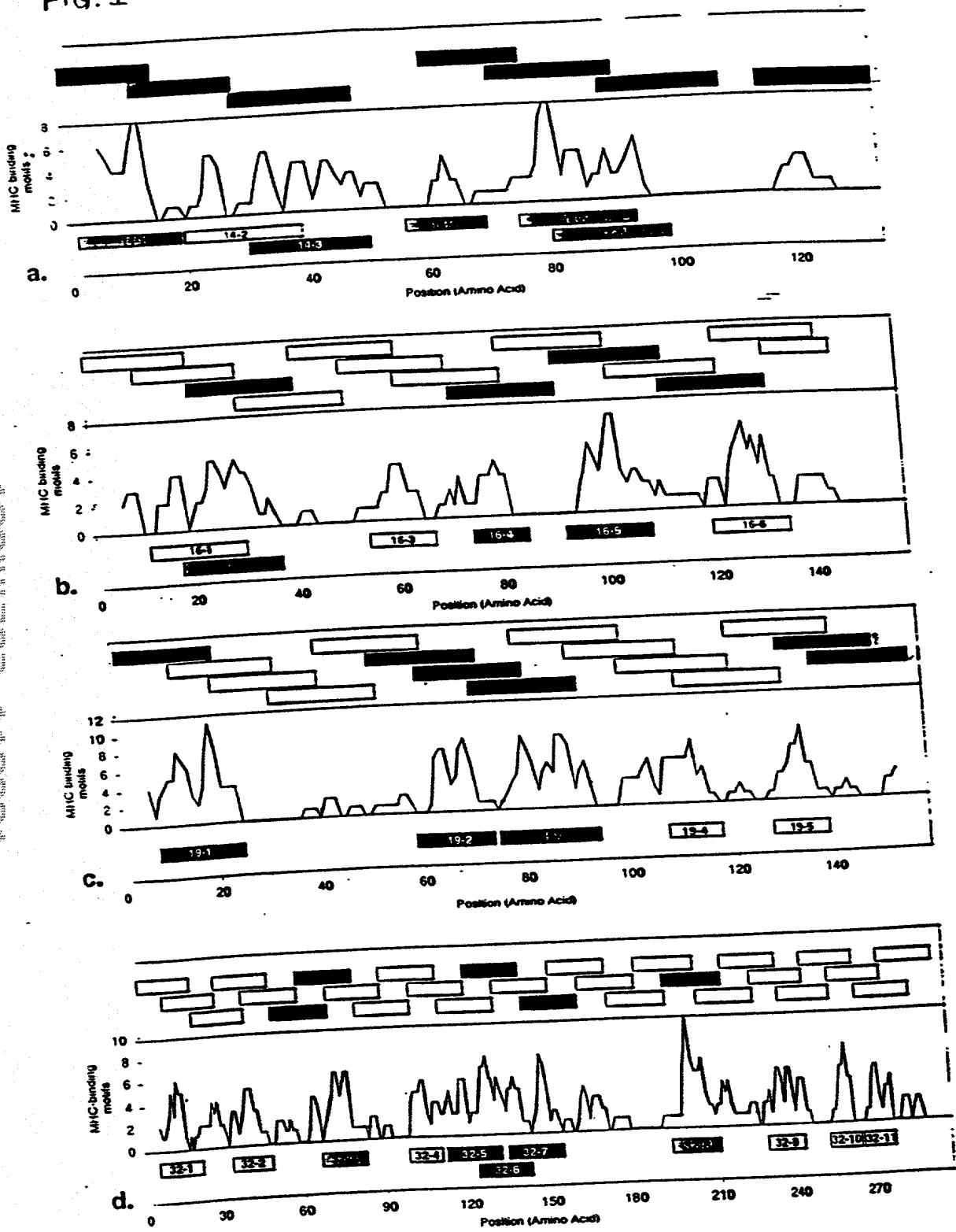


FIG. 2

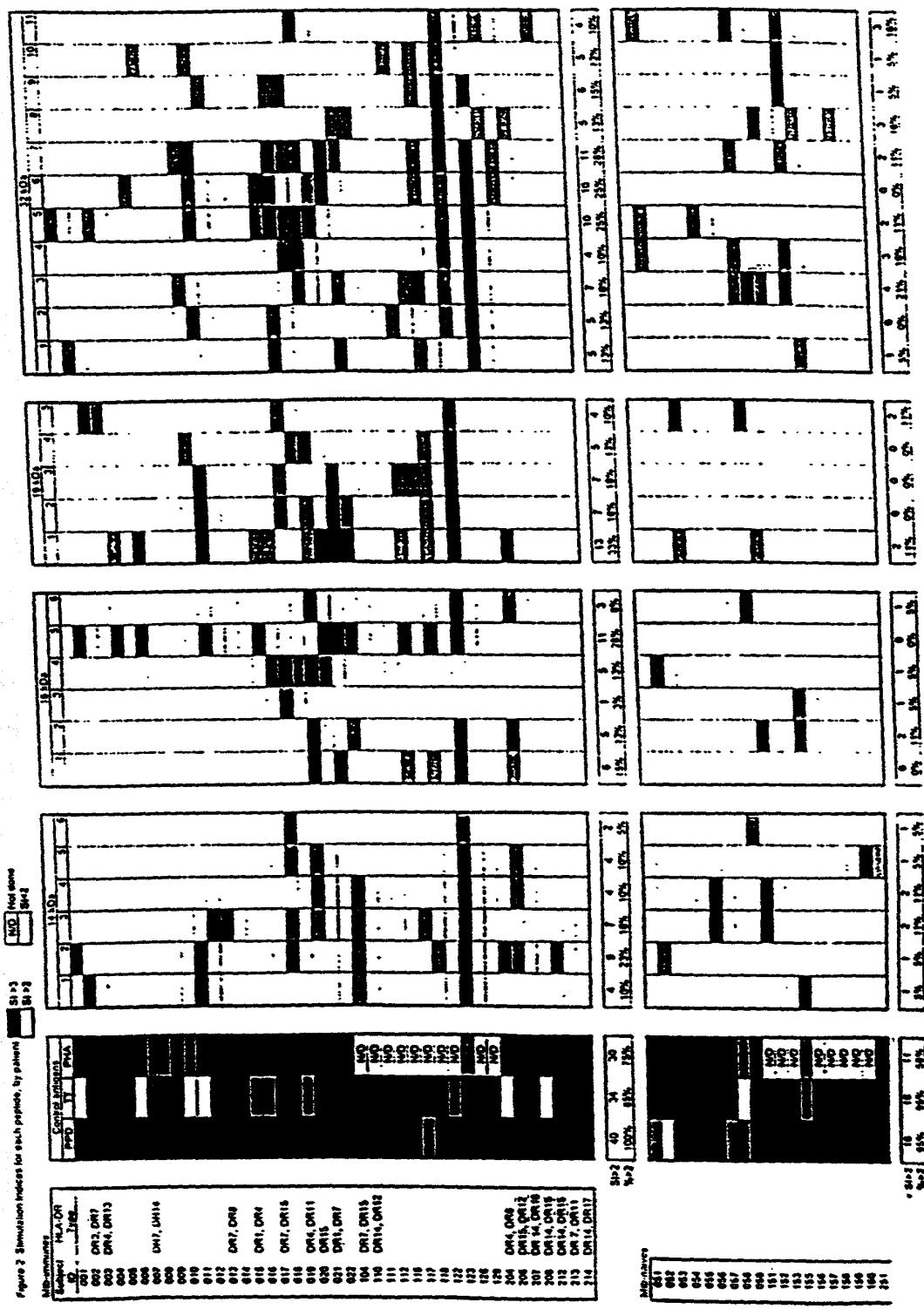


FIG. 3

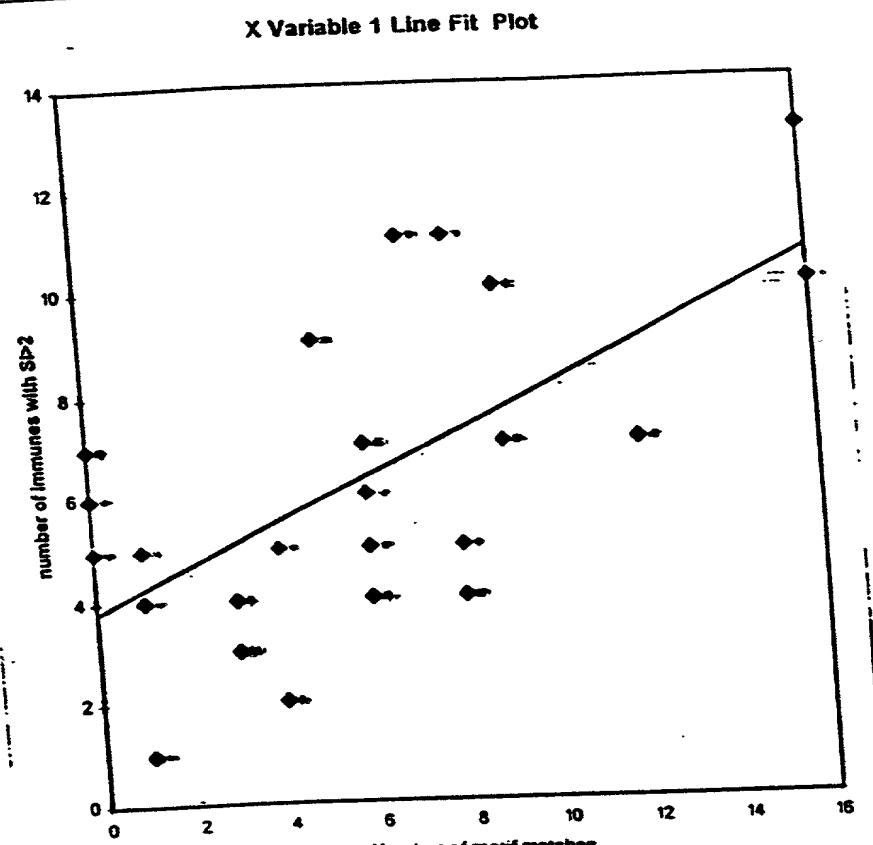


FIG. 4

Name	Start	Stop	Sequence	L	mm	Study	plate label	Prov	Phil	Gambia	combined
14-1	1	18	MLGNARSVPNTTLGMHC (SEQ ID NO:1)	18	14	Brian	14-1	9%	10%		10%
14-2	19	38	GSFOSAPSNQWLKLGLVERFG (SEQ ID NO:2)	20	14	Brian	14-2	14%	18%		17%
14-3	30	49	LKLGLVERGVAKLNAEVMS (SEQ ID NO:3)	20	16	Brian	14-3	18%	13%		15%
14-4	56	68	QAVMLGTTGPNRA (SEQ ID NO:4)	13	3	Brian	14-4	5%	13%		10%
14-5	74	93	CEVWSNVSETISGPRLYGEM (SEQ ID NO:5)	20	10	Brian	14-5	9%	13%		12%
14-6	80	99	VSETISGPRLYGEMTMQGTR (SEQ ID NO:6)	20	8	Brian	14-6	5%	5%		5%
16-1	11	30	RSLFPEFSELFAAFPSFAGL (SEQ ID NO:7)	20	12	Brian	16-1	9%	15%		13%
16-2	17	36	FSELFAAFPSFAGLRLPTFDT (SEQ ID NO:8)	20	12	Brian	16-2	5%	15%		11%
16-3	54	66	AELPGVDPDKDVD (SEQ ID NO:9)	13	2	Brian	16-3	5%	3%		4%
16-4	74	84	LTIKAERTEQK (SEQ ID NO:10) FAYGSFVRTVSLPVGAA (SEQ ID NO:11)	11	1	Brian	16-4	23%	0%		8%
16-5	92	107	16	10	Brian	16-5	32%	18%		13%	
16-6	119	135	GILTVSYAYSEGKPTEK (SEQ ID NO:12)	17	9	Brian	16-6	5%	10%		8%
19-1	7	25	VAVAGAAILVAGLSGCSSN (SEQ ID NO:13)	19	23	Brian	19-1	36%	23%		28%
19-2	58	74	QNVIGSVVCTTAAGNNV (SEQ ID NO:14)	17	10	Brian	19-2	18%	8%		12%
19-3	75	94	IAGGAATGAAVLTDGNPP (SEQ ID NO:15)	20	19	Brian	19-3	14%	15%		15%
19-4	107	117	VTLGTYSGTQQ (SEQ ID NO:16)	11	3	Brian	19-4	14%	10%		11%
19-5	127	139	SHYKITGJATGVD (SEQ ID NO:17)	13	5	Brian	19-5	14%	8%		10%
32-1	6	20	LPVEYLQVSPSMGR (SEQ ID NO:18)	15	9	Brian	32-1	9%	10%		10%
32-10	252	264	GVFDPPDSGTHSW (SEQ ID NO:27)	13	5	Brian	32-10	9%	15%		13%
32-11	265	276	EYWGQLNAMKP (SEQ ID NO:28)	12	4	Brian	32-11	5%	15%		11%
32-2	35	47	ALYILDGLRAQDD (SEQ ID	13	4	Brian	32-2	9%	8%		8%

38G-19	346	358	ASFLDQVHFQQLP (SEQ ID NO:45)	13	2	Krista	38G-19	0%	24%	17%
38KD-1	3	22	IRLHTLLAVLTAAPLLAAG (SEQ ID NO:46)	20	23	Rose	38-1	60%	18%	17%
38KD-10	272	291	NFLIPDAQSIQAAAAGFASK (SEQ ID NO:47)	20	16	Rose	38-3	50%	18%	22%
38KD-11	316	335	YAIVNVRQDAATAQTLQAF (SEQ ID NO:48)	20	10	Rose	38-6	40%	18%	22%
38KD-4	86	102	TGSGAGIAQAAAGTVNI (SEQ ID NO:49)	17	10	Rose	38-5	45%	18%	28%
38KD-5	114	133	MAAHKGIMNIALAISAAQVN (SEQ ID NO:50)	20	16	Rose	38-2	45%	27%	17%
38KD-7	169	188	LNPQGVNLPGTAVVPLHRSQG (SEQ ID NO:51)	20	10	Rose	38-4	10%	0%	22%
38KD-9	239	258	AETPGCVAYIGISFLDQASQ (SEQ ID NO:52)	20	9	Rose	38-7	25%	55%	17%
38U-01	3	23	IRLHTLLAVLTAAPLLAAG (SEQ ID NO:53)	19	23	Krista	38U-01	0%	28%	28%
38U-02	37	49	AGAGITVATTPASS (SEQ ID NO:54)	13	5	Krista	38U-02	23%	17%	19%
38U-05	86	103	TGSGAGIAQAAAGTVNIG (SEQ ID NO:55)	18	10	Krista	38U-05	0%	28%	28%
38U-06	113	135	DMAAHKGIMNIALAISAAQVN (YN (SEQ ID NO:36))	23	16	Krista	38U-06	15%	24%	21%
38U-09	164	177	PQIAALNPQVNLPQG (SEQ ID NO:57)	14	5	Krista	38U-09	31%	14%	19%
38U-14	273	295	FLLPDAQSIQAAAAGFASKTPA (N (SEQ ID NO:58))	23	17	Krista	38U-14	23%	10%	14%
E6G-01	7	19	NFAGIEAAASAIQ (SEQ ID NO:59)	13	6	Krista	E6G-01	38%	50%	46%
E6U-01	5	24	QWNFAGIEAAASAIQGNVTS (SEQ ID NO:60)	20	9	Krista	E6U-01	38%	32%	34%
ESAT6- 1a	7	26	NFAGIEAAASAIQGNVTSIH (SEQ ID NO:61)	20	10	Rose	E6-1	45%	9%	17%
ESAT6- 1b	18	37	IQGNVNTSIIHLDEGKQSLT (SEQ ID NO:62)	20	7	Rose	E6-2	25%	36%	17%
ESAT6- 2	63	78	TELNNALQNLARTISE (SEQ ID NO:63)	16	8	Rose	E6-3	85%	27%	6%
MPT59- 1	15	34	LMIGTAAAVVLPGLVGLGG (SEQ ID NO:64)	20	21	Rose	59-1	35%	27%	33%
MPT59- 10	305	324	EYWGAQLNAMKGDLQSSLGA (SEQ ID NO:65)	20	8	Rose	59-5	35%	9%	6%
MPT59- 3	100	115	WYYQSGLSTVMPVGGQ (SEQ ID NO:66)	16	11	Rose	59-3	35%	0%	17%
MPT59- 5	163	178	IGLSMAGSSAMILAAY (SEQ ID NO:81)	16	10	Rose	59-4	35%	36%	11%

MPT59-6	181	199	QQFIYAGSLSALLDPSQGM (SEQ ID NO:67)	19	12	Rose	59-2	40%	9%	11%	21%
MPT63-1a	21	40	VVLGKVKVSDLKSSTAIVPGY (SEQ ID NO:68)	20	10	Rose	63-2	40%	18%	33%	31%
MPT63-1b	27	46	VSDIKSSTAIVPGYPVAGQV (SEQ ID NO:69)	20	10	Rose	63-3	70%	18%	11%	35%
MPT63-2	56	75	TRGSVTPAVSQNARTADGI (SEQ ID NO:70)	20	10	Rose	63-1	40%	27%	11%	26%
MPT64-1	3	22	IKIFMLVTAIVVLLCCSGVAT (SEQ ID NO:71)	20	23	Rose	64-1	35%	27%	11%	25%
MPT64-2a	50	69	YNINISLPSYYPDQKSLENY (SEQ ID NO:72)	20	7	Rose	64+3	70%	36%	6%	39%
MPT64-3	103	122	AIPPRGTQAVVLKVYQNAAGG (SEQ ID NO:73)	20	10	Rose	64-2	60%	64%	11%	44%
MPT70-1a	9	28	ATSFAAAAGLAAALAVAVSPA (SEQ ID NO:74)	20	26	Rose	70-1	30%	27%	28%	28%
MPT70-1b	11	30	SFAAAGLAAALAVAVSPA (SEQ ID NO:75)	20	26	Rose	70-2	55%	36%	11%	35%
MPT70-4	85	97	VNLVDTLNSGQYT (SEQ ID NO:76)	13	7	Rose	70-7	55%	9%	11%	27%
MPT70-5a	104	123	AAFSKLPASTIDELKTNSSL (SEQ ID NO:77)	20	10	Rose	70-5	50%	18%	22%	31%
MPT70-5b	114	133	IDELKTNSSSLTSILTYHVV (SEQ ID NO:78)	20	14	Rose	70-3	85%	64%	22%	58%
MPT70-5c	124	143	LTSILTYHVVAGQTSPANVV (SEQ ID NO:79)	20	13	Rose	70-4	50%	18%	11%	28%
MPT70-6	174	192	GVSTANATVYMDSVLMP (SEQ ID NO:80)	19	11	Rose	70-6	55%	45%	17%	39%

Brian's peptides in Phil = pop. size of 39
 Brian's peptides in Prov = pop size of 22

Rose's peptides in Phil = pop size of 11
 Rose's peptides in Prov = pop size of 20
 Rose's peptides in Gambia = pop size of 18

Krista's peptides in Gambia 1995 = pop size of 12
 Krista's peptides in Gambia 1996 = pop size of 17
 Krista's peptides in Providence 1997 = pop size of 13
 combined = (%resp pop A x pop size A + %resp pop B x popsize B)/(popsize A + popsize B))

L = length
 mm = motif match